



Supplemental_Fig_S14: Unsupervised clustering of Type 2 diabetic samples when excluding marker genes demonstrate distinct clustering by cell type.

Circular dendrogram of Type 2 diabetic single cell samples continues to demonstrate clear separation of cell transcriptomes by cell type. 1900 highly expressed genes with $\log_2(\text{CPM}) > 10.5$ were used in the analysis. Color labels are the same as shown in Figure 3B.